

18 FEB 2003

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re the Application of )  
 )  
Caroline Dean et al. )  
 )  
Serial No. 10/088,187 )  
 )  
Filed: March 15, 2002 )  
 )  
For: "Methods and Means for )  
Modification of Plant )  
Flowering Characteristics")

RE-SUBMISSION OF SEQUENCE LISTING  
UNDER 37 C.F.R. §§1.821-1.825


To comply with the requirements under 37 C.F.R. §§1.821-1.825, re-submitted herewith is a listing of the amino acid and nucleotide sequences presented in the above-referenced application. The sequence listing is being re-submitted in computer-readable form to replace the unreadable original diskette. Applicants respectfully request entry of the sequence listing into the above identified patent application. The undersigned hereby verifies that the computer readable form of the sequence listing is identical to the originally submitted paper copy form and does not contain any new matter.

In the event that a fee is required, the Commissioner is authorized to charge the account of the undersigned, Account No. 04-1406. A duplicate copy of this sheet is enclosed.

Respectfully submitted,

DANN, DORFMAN, HERRELL AND SKILLMAN  
A Professional Corporation

By

  
Kathleen D. Rigaut, Ph.D., J.D.  
PTO Registration No. 43,047

Telephone: (215) 563-4100

## SEQUENCE LISTING

<110> Dean, Caroline  
Levy, Yaron Y

<120> Methods and Means for Modification of Plant Flowering  
Characteristics

<130> 0380-P02825US0

<140> US 10/088,187

<141> 2002-03-15

<150> PCT/GB00/03525

<151> 2000-09-13

<150> GB 9922071.7

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<170> PatentIn Ver. 2.1

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<210> 13

<211> 1494

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: vrnl-2  
mutation

<400> 13

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<210> 14  
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<220>  
 <223> Description of Artificial Sequence: Primer

<400> 14  
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<210> 15  
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<220>  
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<400> 15  
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<210> 16  
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<220>  
 <223> Description of Artificial Sequence: Primer

<400> 16  
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<210> 17  
 <211> 18  
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<220>  
 <223> Description of Artificial Sequence: Primer

<400> 17  
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<210> 18  
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<220>  
 <223> Description of Artificial Sequence: Primer

<400> 18  
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<210> 19  
<211> 20  
<212> DNA  
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<220>  
<223> Description of Artificial Sequence: Primer

<400> 19  
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<210> 20  
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<220>  
<223> Description of Artificial Sequence:  
Oligonucleotide

<400> 20  
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17

<210> 21  
<211> 17  
<212> DNA  
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<220>  
<223> Description of Artificial Sequence:  
Oligonucleotide

<400> 21  
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<210> 22  
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<212> DNA  
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<220>  
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Oligonucleotide

<400> 22  
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19

<210> 23  
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Oligonucleotide

<400> 23  
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<210> 24  
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<212> DNA  
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<220>  
<223> Description of Artificial Sequence:  
Oligonucleotide

<400> 24  
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<210> 25  
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<212> DNA  
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<220>  
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<400> 25  
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<210> 26  
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<220>  
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<400> 26  
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<210> 27  
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<212> DNA  
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<400> 27  
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<210> 28  
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<400> 28  
tttcttgat agtggatgag

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<210> 29  
<211> 21  
<212> DNA  
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<400> 29  
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21

<210> 30  
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<210> 31  
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<400> 31  
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18

<210> 32  
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<220>  
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<400> 32  
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18

<210> 33  
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Oligonucleotide

<400> 33  
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<210> 34  
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<212> DNA  
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<400> 34  
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<210> 35  
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<212> DNA  
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<220>  
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Oligonucleotide

<400> 35  
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17

<210> 36  
<211> 18  
<212> DNA  
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Oligonucleotide

<400> 36  
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18

<210> 37  
<211> 17  
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Oligonucleotide

<400> 37  
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<210> 38  
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<212> DNA  
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Oligonucleotide

<400> 38  
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<210> 39  
<211> 18  
<212> DNA  
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Oligonucleotide

<400> 39  
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18

<210> 40  
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<400> 40  
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<210> 41  
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<400> 41  
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18

<210> 42  
<211> 17  
<212> DNA  
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<220>  
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Oligonucleotide

<400> 42  
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17

<210> 43  
<211> 19  
<212> DNA  
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Oligonucleotide

<400> 43  
ccttctgttt ctgtttctc

19

<210> 44  
<211> 19  
<212> DNA  
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<220>  
<223> Description of Artificial Sequence:  
Oligonucleotide

<400> 44  
gagaaacaga aacagaagg

19

<210> 45  
<211> 18  
<212> DNA  
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<400> 45  
aagatactcc tacacgac

18



<210> 46  
 <211> 19  
 <212> DNA  
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<220>  
 <223> Description of Artificial Sequence:  
 Oligonucleotide

<400> 46  
 gtctcgtttt ttctctcgg

19

<210> 47  
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 <212> DNA  
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<400> 47  
 ctaccacagt tcccacctac

20

<210> 48  
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 <212> PRT  
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 20 25 30  
 Asp Gly Trp Gln Glu Phe Val Asn Arg Phe Ser Ile Arg Ile Gly Phe  
 35 40 45  
 Arg Tyr Lys Val Thr Val Tyr Ile Phe Gln Phe Tyr Pro Pro His Ser  
 50 55 60  
 Glu Ile Asn His His Ser Ser Ser Glu Ala Leu Met Gln Met Asp Ser  
 65 70 75 80  
 Ala Gln Asn Gln Phe Asn Lys Arg Ala Arg Leu Phe Glu Asp Pro Glu  
 85 90 95  
 Leu Lys Asp Ala Lys Val Ile Tyr Pro Ser Asn Pro Glu Ser Thr Glu  
 100 105 110  
 Pro Val Asn Lys Gly Tyr Gly Gly Ser Thr Ala Ile Gln Ser Phe Phe  
 115 120 125  
 Lys Glu Ser Lys Ala Glu Glu Thr Pro Lys Val Leu Lys Lys Arg Gly  
 130 135 140

Arg	Lys	Lys	Lys	Asn	Pro	Asn	Pro	Glu	Glu	Val	Asn	Ser	Ser	Thr	Pro	145	150	155	160
Gly	Gly	Asp	Asp	Ser	Glu	Asn	Arg	Ser	Lys	Phe	Tyr	Glu	Ser	Ala	Ser	165	170		175
Ala	Arg	Lys	Arg	Thr	Val	Thr	Ala	Glu	Glu	Arg	Glu	Arg	Ala	Val	Asn	180	185		190
Ala	Ala	Lys	Thr	Phe	Glu	Pro	Thr	Asn	Pro	Tyr	Phe	Arg	Val	Val	Leu	195	200		205
Arg	Pro	Ser	Tyr	Leu	Tyr	Arg	Gly	Cys	Ile	Met	Tyr	Leu	Pro	Ser	Gly	210	215		220
Phe	Ala	Glu	Lys	Tyr	Leu	Ser	Gly	Ile	Ser	Gly	Phe	Ile	Lys	Leu	Gln	225	230		235
Leu	Gly	Glu	Lys	Gln	Trp	Pro	Val	Arg	Cys	Leu	Tyr	Lys	Ala	Gly	Arg	245	250		255
Ala	Lys	Phe	Ser	Gln	Gly	Trp	Tyr	Glu	Phe	Thr	Leu	Glu	Asn	Asn	Ile	260	265		270
Gly	Glu	Gly	Asp	Val	Cys	Val	Phe	Glu	Leu	Leu	Arg	Thr	Arg	Asp	Phe	275	280		285
Val	Leu	Glu	Val	Thr	Ala	Phe	Arg	Val	Asn	Glu	Tyr	Val				290	295		300